



Multimodal radiomics based on fluorine-18 prostate-specific membrane antigen positron emission tomography and multiparametric magnetic resonance imaging in predicting persistent prostate-specific antigen after radical prostatectomy

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Background: Persistent prostate-specific antigen (PSA) after radical prostatectomy (RP) is associated with increased metastasis and mortality. However, the value of the radiomics for predicting persistent PSA is unclear. Our study aimed to evaluate the diagnostic performance of ¹⁸F-PSMA-1007 positron emission tomography (PET) and multiparametric magnetic resonance imaging (mpMRI) radiomics for the prediction of persistent PSA after RP.

Methods: Retrospective analysis was performed on 141 patients with prostate cancer (PCa) who had undergone ¹⁸F-prostate-specific membrane antigen (PSMA)-1007 PET and mpMRI scans before RP. Patients were placed into two groups according to PSA levels examined within 4–8 weeks after surgery: a nonpersistent PSA group and a persistent PSA group. PET-derived and mpMRI-derived radiomics features were used to develop radiomics models. Age and initial PSA were incorporated into the clinical model. Individual models and their various combinations were developed and their performance evaluated.

Results: All radiomics models consistently outperformed the clinical model [C model: area under curve (AUC) =0.744]. The best-performing radiomics model was the PET- and mpMRI-derived model (PM model) created by combining the radiomics features of PET and mpMRI, which yielded an AUC of 0.849 in the validation cohort, and was superior to the other radiomics models, including the PET-derived model (P model: AUC =0.794) and the mpMRI-derived model (M model: AUC =0.815). The combined model, integrating the clinical variables and the best-performing radiomics model, demonstrated the highest performance (AUC =0.903) and significantly outperformed the C model (P<0.05). Decision curve analysis indicated that the combined model provided greater net benefits than did the C model and PM model.

Conclusions: The combined radiomics-clinical model was the best-performing model and outperformed both clinical and radiomics models in predicting persistent PSA, indicating that clinical variables can complement PSMA-PET and mpMRI radiomics for early risk stratification following RP.

Keywords: Prostate cancer (PCa); persistent prostate-specific antigen (persistent PSA); PSMA-1007 positron emission tomography (PET); multiparametric magnetic resonance imaging (mpMRI); radiomics

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Introduction

For patients with localized and locally progressed prostate cancer (PCa), radical prostatectomy (RP) provides favorable long-term oncological outcomes (1-3). However, a significant number of patients may experience disease recurrence after RP. Prostate-specific antigen (PSA) measurements play a crucial role in patient follow-up. The guidelines from the European Association of Urology (EAU) recommend a PSA assessment within 3 months after RP (4). Given the short half-life of PSA, it is theoretically possible that PSA is at an undetectable low level within 4–8 weeks after surgery. However, PSA can still be detected in 5% to 30% of patients with PCa after RP due to the presence of undetected micrometastases, residual malignant lesions, and residual benign tissue at the surgical margin (5). Persistent PSA is generally defined as a PSA level ≥ 0.1 ng/mL within 4–8 weeks after RP (6-15). Previous studies have discovered that persistent PSA is a powerful prognostic predictor for the development of metastasis and death following RP (6,8,12,15). Therefore, accurate prediction of persistent PSA before patients experience biochemical recurrence (BCR) could help identify high-risk patients with poor tumor prognosis in clinical practice, which could further assist physicians in determining which individuals are most likely to benefit from follow-up and adjuvant therapy.

In clinical practice, multiparametric magnetic resonance imaging (mpMRI) serves as the primary imaging modality to evaluate PCa. Many retrospective studies have highlighted the unique advantages of mpMRI in assessing PCa (16-18). Recently, positron emission tomography (PET) with prostate-specific membrane antigen (PSMA) targeting has come into prominence as a novel imaging evaluation approach for PCa, due to its whole-body imaging capabilities and the advantages of specific prostate targeting (19-22). ^{18}F -labeled PSMA-targeted radiopharmaceuticals have a higher image resolution compared to other radiopharmaceuticals (21,23). Notably, in clinical applications, ^{18}F -PSMA-1007 has been successfully applied in diagnosing PCa and has demonstrated good performance (24). The integration of clinical, radiological, and functional imaging parameters before surgery has also been shown to be effective in predicting persistent PSA (25).

As a novel noninvasive technique, radiomics aims to extract and analyze a wide range of quantitative imaging features from radiographic images with high throughput. This facilitates a comprehensive characterization of intratumoral information and changes (26). Handcrafted radiomics features have been used in previous studies to assist with the detection and prognosis of various cancers, including breast cancer (27), pancreatic neuroendocrine tumor (28), and lung cancer (29). Recently, the application of radiomics in PCa has garnered intense extensive research attention due to its potentially significant benefits for healthcare systems (30-33). A number of studies have reported on radiomics improving PCa detection, International Society of Urological Pathology (ISUP) Grade Group prediction, and BCR in patients with primary PCa; however, there is little evidence regarding the use of multimodal radiomics in predicting persistent PSA (31).

The study aimed to determine the value of preoperative ^{18}F -PSMA-1007 PET and mpMRI radiomics in predicting persistent PSA after RP. We hypothesize that radiomics could be used to predict persistent PSA. We present this article in accordance with the TRIPOD reporting checklist (available at <https://qims.amegroups.com/article/view/10.21037/qims-24-2162/rc>).

Methods

Patient population

A total of 321 patients with histopathologically proven localized PCa who underwent RP at The First Affiliated Hospital of Wenzhou Medical University between January 2019 and March 2022 were included in this retrospective analysis. The laparoscopic RP approach was applied to patients including but not limited to those whose preoperative examination showed positive lymph nodes and who underwent lymph node dissection. The exclusion criteria were as follows: (I) adjuvant therapy administered before imaging exam or operation, (II) adjuvant therapy administered within 8 weeks after surgery, (III) lack of complete clinical and pathological data, and (IV) patients without complete preoperative ^{18}F -PSMA-1007 PET and mpMRI data. The final analysis included a cohort

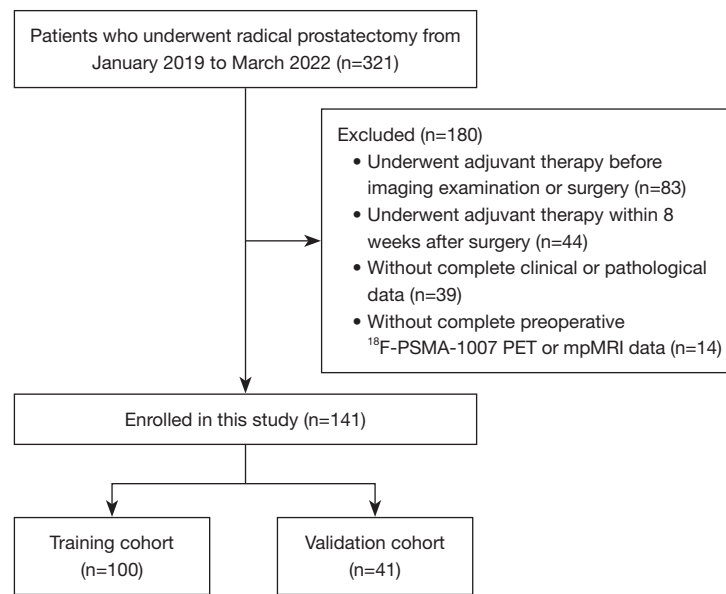


Figure 1 Flowchart of the patient selection process. mpMRI, multiparametric magnetic resonance imaging; PSMA, prostate-specific membrane antigen; PET, positron emission tomography.

of 141 patients. The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013) and was approved by the Ethics Committee of The First Affiliated Hospital of Wenzhou Medical University (No. KY2022-R012). Informed consent was waived due to the retrospective nature of the analysis. The patient selection process is visually represented in *Figure 1*.

Imaging protocol

For each patient involved in the study, a hybrid PET-computed tomography (CT) scanner (Gemini TF 64, Philips Healthcare, Best, the Netherlands) was used to obtain the ^{18}F -PSMA-1007 PET images. Additionally, 3.0-T MR scanners (Signa HDxt \times t3.0 T, GE Healthcare, Chicago, IL, USA; Achieva 3.0 T, Philips Healthcare) were used to conduct the mpMRI experiments. The mpMRI scans comprised a T2-weighted (T2W) sequence, a diffusion-weighted imaging (DWI) sequence, and apparent diffusion coefficient (ADC) maps. The details are provided in [Appendix 1](#).

Image segmentation and radiomics feature extraction

All procedures were carried out in accordance with the Image Biomarker Standardization Initiative principles (34).

The workflow for feature extraction is illustrated in *Figure 2*. To extract the radiomic features, the volumes of interest (VOIs) were first manually segmented in both ^{18}F -PSMA-1007 PET and mpMRI images.

For the PET images, LIFEx software (version 6.30; <https://www.lifexsoft.org>) was used to segment the whole prostate. During the segmentation process, PET VOIs were defined based on (I) the whole prostate ($\text{VOI}_{\text{whole}}$) and (II) the threshold at 40% of standardized uptake value (SUV_{max}) ($\text{VOI}_{40\%}$). A previous study reported that at a 40% SUV_{max} , radiomics features based on the ^{18}F -PSMA-1007 PET demonstrate optimal predictive performance in assessing the extracapsular extension (ECE) of PCa (35). Before feature extraction, all VOIs underwent normalization and discretization with a fixed bin width (default = 0.3) and subsequently were resampled to $2 \times 2 \times 2 \text{ mm}^3$. Radiomics features were extracted from the defined VOIs ($\text{VOI}_{\text{whole}}$ and $\text{VOI}_{40\%}$) in PET images using LIFEx software. For each of the defined VOIs, conventional features (SUV_{min} , SUV_{max} , SUV_{mean} , etc.), shape features, and texture features were extracted separately.

For the mpMRI images, we used ITK-SNAP software (version 3.6.0; <http://www.itksnap.org>) to manually segment the whole prostate on DWI and T2W images in a slice-by-slice approach. Subsequently, rigid registration was performed to align the VOIs delineated on DWI scans with

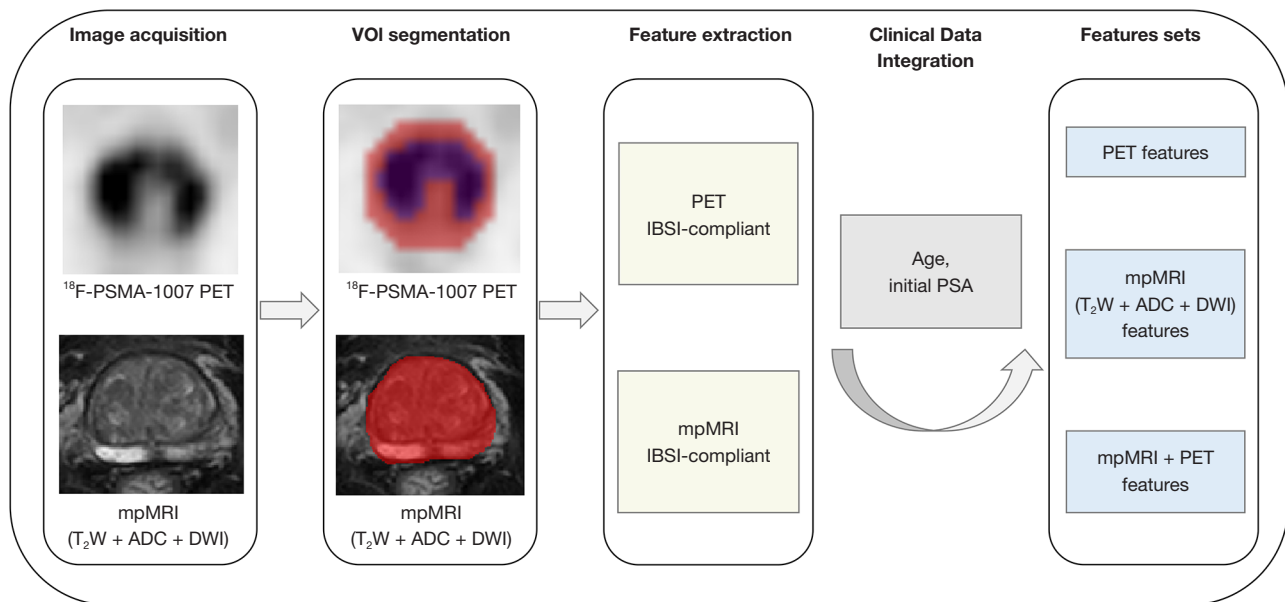


Figure 2 Feature extraction workflow for ^{18}F -PSMA-1007 PET and mpMRI images. ADC, apparent diffusion coefficient; DWI, diffusion-weighted imaging; IBSI, Image Biomarker Standardization Initiative; mpMRI, multiparametric magnetic resonance imaging; PET, positron emission tomography; PSA, prostate-specific antigen; PSMA, prostate-specific membrane antigen; T2W, T2-weighted; VOI, volume of interest.

the corresponding regions on ADC maps, via ITK-SNAP software. Since ADC maps are derived from DWI data, rigid registration was sufficient to ensure spatial consistency. Finally, radiomics features extraction from mpMRI was carried out using the “Pyradiomics” package (version 2.2.0) in Python software (Python Software Foundation, Wilmington, DE, USA). A total of 1,316 features were extracted from the VOIs of each image sequence (T_2W , DWI, and ADC), which included seven categories: (I) first order, (II) shape features, and texture features derived from a (III) gray-level co-occurrence matrix (GLCM), (IV) gray-level difference method (GLDM), (V) gray-level run-length matrix (GLRLM), (VI) gray-level size zone matrix (GLSZM), and (VII) neighborhood gray-tone difference matrix (NGTDM).

The segmentations were carried out independently by two nuclear medicine physicians (with 3 and 5 years of experience interpreting PSMA hybrid imaging, respectively) and two radiologists (with 3 and 5 years of experience reading prostate mpMRI). To guarantee impartial and unbiased results, all doctors were kept blind to patient-specific information during the VOIs delineation. The intraclass correlation coefficient (ICC) with one-way analysis of variance (ANOVA) was used to assess

interobserver reproducibility. High feature stability was indicated by an ICC of ≥ 0.75 . All parameters with an ICC < 0.75 were initially eliminated in order to decrease unstable features.

Prediction models

First, the preserved stable radiomics features were sorted using the minimum redundancy maximum relevance (mRMR) algorithm, with the top 15 radiomic features most related to persistent PSA being selected for each model. Subsequently, the least absolute shrinkage and selection operator (LASSO) algorithm with 10-fold cross-validation was used to select the most optimized feature subsets for building the radiomics models. To identify persistent PSA, multivariate logistic regression was used to develop prediction models using the previously selected optimized feature subsets on the training cohort. The prediction capacity of models was then computed on the validation cohort. Different prediction models were established. We first established and compared the predictive performance of two PET radiomics models based on the defined VOIs ($\text{VOI}_{\text{whole}}$ and $\text{VOI}_{40\%}$) and chose the model with better performance as the final PET radiomics model (referred

to as the P model). Additionally, a mpMRI radiomics model (referred to as the M model) based on T2W, DWI, and ADC radiomics features and a multimodal radiomics model (referred to as the PM model) integrating the radiomics features from both the PET and mpMRI were established. Age and initial PSA level were incorporated as variables in the clinical model (referred to as the C model) because these variables have long been considered to be independent predictors in PCa risk calculators (36). Finally, the clinical variables (age + initial PSA) were added to the best-performing radiomics model to further investigate whether a combined radiomics-clinical model (referred to as the combined model) might more accurately distinguish persistent PSA from nonpersistent PSA. The area under curve (AUC), sensitivity, specificity, accuracy, and F1 value were used to assess these models' prediction ability.

As only 26 patients had persistent PSA after RP (~18%), oversampling the less common category (PSA ≥ 0.1) in the training set was addressed by using the synthetic minority oversampling technique (SMOTE) to achieve a 1:1 ratio to the most common category (PSA < 0.1). The prediction models were trained with the augmented training data, and then the ability of the trained models to predict persistent PSA was evaluated using the unaugmented validation data.

Statistical analysis

Statistical analyses were performed using SPSS software (version 26.0; IBM Corp., Armonk, NY, USA), MedCalc software (version 19.8, MedCalc Software, Mariakerke, Belgium), and R software (The R Foundation for Statistical Computing; version 4.1.2; <https://www.R-project.org>). For SPSS, the *t* test or Mann-Whitney test was used to compare the differences between continuous variables, while the Fisher exact test or Chi-squared test was applied for categorical variables. For R software, multivariable logistic regression was performed to create the prediction models, and decision curve analysis (DCA) was conducted to evaluate the clinical applicability of the models. The DeLong test was employed to examine the accuracy of prediction between the different models, and the net reclassification improvement (NRI) method was used to assess the incremental value of the models. MedCalc software was used for the assessment of model performance via AUC values and confusion matrix-derived indices. The reproducibility of VOI segmentation was evaluated using the Dice similarity coefficient (DSC), which quantifies the pairwise overlap between repeated segmentations. The DSC

values were calculated using R software, with values ranging from 0 to 1. Higher values indicated greater overlap, and a DSC value exceeding 0.70 was considered indicative of good segmentation reproducibility. All statistical tests were conducted in two-sided fashion, with statistical significance defined as a *P* value of 0.05 or less.

Results

Study population and baseline characteristics

The study included 141 patients who met the criteria for participation. Based on their PSA levels within 4–8 weeks after RP, they were classified into groups: a persistent PSA (26 patients) group and a nonpersistent PSA group (115 patients). Subsequently, at a 7:3 ratio, the patients were randomly separated into a training cohort (*n*=100) and a validation cohort (*n*=41). *Table 1* provides an overview of the baseline characteristics of these patients. No statistically significant differences were observed in baseline characteristics, such as age and weight, between the training and validation cohorts (*P*>0.05).

Model evaluation and comparison

A summary of the diagnostic performance of the established models for the classification of nonpersistent PSA versus persistent PSA (PSA < 0.1 vs. ≥ 0.1) within 4–8 weeks following RP is provided in *Table 2* and *Figure 3*.

The reproducibility of manual segmentation of VOIs was good (average DSCs: 0.82 for PET, 0.78 for DWI, and 0.80 for T2WI). For the PET-derived radiomics models, the model based on VOI_{40%} performed better than did the VOI_{whole}, and thus we chose the model based on the radiomics features from VOI_{40%} as our P model (*Appendix 1*). Overall, The PM model, a multimodal radiomics model, was the best-performing radiomics model, with the highest AUC among other radiomics models in both the training [AUC =0.893; 95% confidence interval (CI): 0.815–0.946] and validation cohorts (AUC =0.849; 95% CI: 0.72–0.941). The P model yielded in an AUC of 0.819 (95% CI: 0.730–0.889) and 0.794 (95% CI: 0.639–0.904) in the training and validation cohorts. Meanwhile, the AUC value of the M model for predicting patients with persistent PSA was 0.862 (95% CI: 0.779–0.923) in the training cohort and 0.815 (95% CI: 0.663–0.919) in the validation cohort. The C model's AUCs were 0.776 (95% CI: 0.681–0.853) and 0.744 (95% CI: 0.583–0.867)

Table 1 Comparison of baseline characteristics between patients in the training and validation cohorts

Characteristic	Training cohort (n=100)	Validation cohort (n=41)	P value
Age (years), median [IQR]	69 [64–72]	67 [62–73]	0.674
Weight (kg), median [IQR]	69.45 [62–75]	65 [58.25–74]	0.054
Initial PSA (ng/mL), median [IQR]	10.66 [7.48–17.65]	10.66 [7.83–22.80]	0.773
Persistent PSA			0.789
Yes	19 (19%)	7 (17.07%)	
No	81 (81%)	34 (82.93%)	
ISUP GG			0.153
1	5 (5%)	2 (4.9%)	
2	36 (36%)	11 (26.8%)	
3	39 (39%)	13 (31.7%)	
4	10 (10%)	4 (9.8%)	
5	10 (10%)	11 (26.8%)	
pT stage			0.885
T2	52 (52%)	20 (48.8%)	
T3a	27 (27%)	11 (26.8%)	
T3b	17 (17%)	9 (22%)	
T4	4 (4%)	1 (2.4%)	
pN stage			0.822
0	95 (95%)	40 (97.6%)	
1	5 (5%)	1 (2.4%)	
ECE			0.884
Yes	45 (45%)	19 (46.3%)	
No	55 (55%)	22 (54.7%)	
PSM			0.144
Yes	31 (31%)	18 (44%)	
No	69 (69%)	23 (56%)	
VI			0.629
Yes	18 (18%)	6 (14.6%)	
No	82 (82%)	35 (85.4%)	

ECE, extracapsular extension; IQR, interquartile range; ISUP GG, International Society of Urological Pathology Group Grade; PSA, prostate-specific antigen; PSM, positive surgical margin; VI, vascular invasion.

in the training and validation cohorts, respectively. When all radiomics models were compared to the C model alone, all demonstrated superior performance. The Delong test indicated no significant difference in AUC values between any of the radiomics models (P model, M model, and PM

model) and the C model ($P>0.05$). However, the NRI indicated that the radiomics models, except the P model, could improve the ability to predict persistent PSA as compared to the C model ($P<0.05$),

The combined model constructed by combining the

Table 2 Model performance

Models	Training cohort					Validation cohort				
	AUC	ACC	SEN	SPE	F1	AUC	ACC	SEN	SPE	F1
C model	0.776	0.780	0.632	0.815	0.522	0.744	0.756	0.714	0.765	0.500
M model	0.862	0.780	0.842	0.765	0.593	0.815	0.707	0.714	0.706	0.455
P model	0.819	0.810	0.737	0.827	0.596	0.794	0.707	0.714	0.706	0.455
PM model	0.893	0.820	0.842	0.815	0.640	0.849	0.756	0.857	0.735	0.546
Combined	0.914	0.830	0.842	0.827	0.653	0.903	0.780	0.857	0.765	0.571

AUC, area under the curve; ACC, accuracy; C model, clinical model; M model, multiparametric magnetic resonance imaging-derived model; P model, positron emission tomography-derived model; PM model, PET- and mpMRI-derived model; SEN, sensitivity; SPE, specificity.

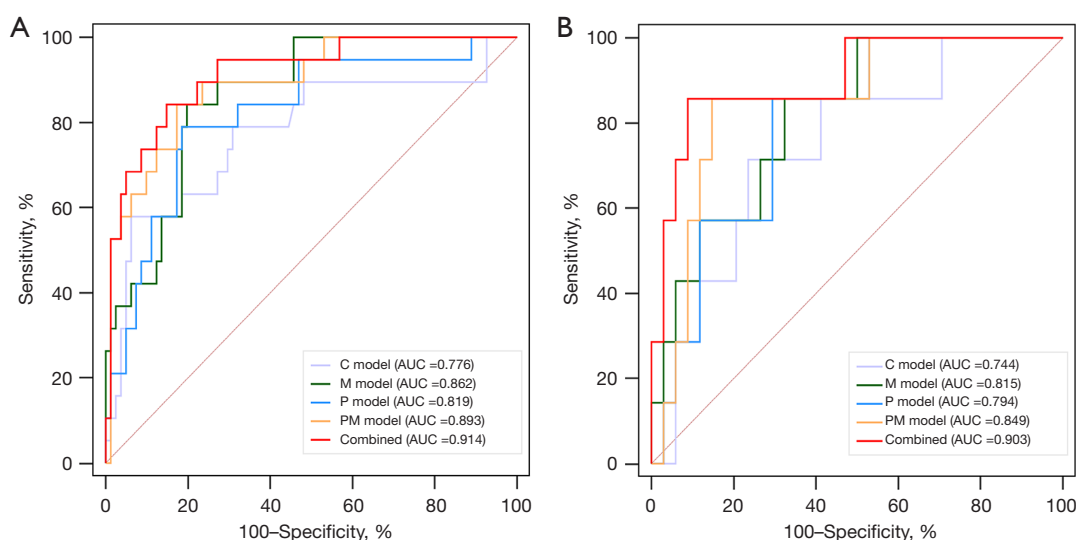


Figure 3 The ROC curve analyses of the different models for predicting persistent PSA in the (A) training cohort and (B) validation cohort, respectively. AUC, area under the curve; C model, clinical model; M model, multiparametric magnetic resonance imaging-derived model; mpMRI, multiparametric magnetic resonance imaging; P model, positron emission tomography-derived model; PET, positron emission tomography; PM model, PET- and mpMRI-derived model; PSA, prostate-specific antigen; ROC, receiver operating characteristic.

C model with the best-performing radiomics model was the best model, with AUCs in the training and validation cohorts of 0.914 (95% CI: 0.840–0.960) and 0.903 (95% CI: 0.770–0.973), respectively. These findings highlight the value of combining radiomics features with clinical variables to improve the accuracy of predicting persistent PSA after RP. According to the DeLong test, the combined model and the C model had statistically significantly different AUC values ($P < 0.05$). Additionally, DCA indicated that as compared to the C model or PM model and a process with all or no intervention, the combined model yielded larger net benefits within a given threshold probability (details

in Figure 4). The details of the radiomic features and the parameters of the logistic regression model are shown in Appendix 2.

Discussion

In this study, we conducted an analysis of patients who underwent RP to assess the diagnostic accuracy of radiomics based on preoperative mpMRI and ^{18}F -PSMA-1007 PET in the identification of persistent PSA. Overall, our analysis revealed that the combined model included both clinical variables and the best-performing radiomics model yielded

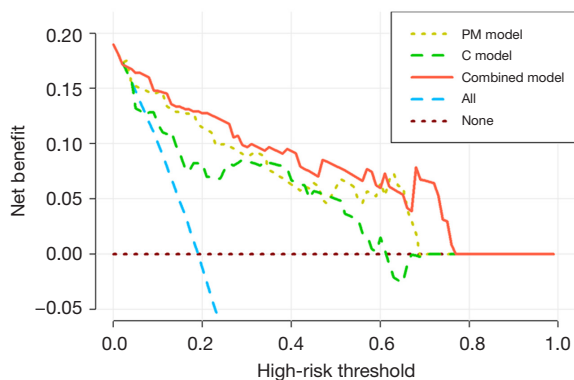


Figure 4 Decision curves of the PM model, C model, and the combined model for predicting persistent PSA in the validation cohort. C model, clinical model; PM model, positron emission tomography- and multiparametric magnetic resonance imaging-derived model; PSA, prostate-specific antigen.

a good predictive performance and outperformed all the radiomics models (P model, M model, and PM model), as well as the clinical baseline model (C model).

Identifying early persistent PSA after RP has significant clinical implications, as it is strongly correlated with poor metastasis-free survival, cancer-specific survival, and overall survival (6). Patients with persistent PSA tend to have more aggressive tumors and poorer oncology outcomes compared to those with undetectable PSA levels after RP (37). Therefore, accurately identifying those patients with persistent PSA after RP can assist in providing personalized treatment options so that patients may benefit from further treatment. Although previous research has focused on pathologic features and conventional imaging parameters for risk stratification in patients with persistent PSA (25), the value of radiomics in this context remains unclear. Our study represents the first comprehensive assessment of the potential of preoperative mpMRI and ^{18}F -PSMA-1007 PET-based radiomics for predicting persistent PSA after RP. Consequently, our findings may offer novel insights and valuable contributions to this area of research.

The importance of hybrid PSMA PET and mpMRI in diagnosing, staging, and identifying recurrent lesions in PCa has been the subject of a thorough investigation (38-41). Its capacity to detect primary tumors and local recurrence, along with its utility in assessing capsular invasion and lymph node staging within the prostate gland, has been substantiated (42,43). Notably, some studies have suggested that this hybrid imaging technique may be more effective than MRI or PET alone in detecting PCa (44-46).

Radiomics analysis has been widely employed in patients diagnosed with primary PCa (31). A significant number of these studies have primarily used MRI and have effectively demonstrated the ability of radiomics to accurately characterize and detect clinically significant PCa, as well as provide valuable insights into ECE and the prediction of BCR (32,33). Limited evidence exists regarding the clinical significance of combining PET and MRI radiomics features in predicting PCa. Basso Dias *et al.* conducted a study of patients with histopathologically confirmed PCa lesions with different ISUP grades to assess the diagnostic accuracy of radiomics features for identifying medium or high-risk tumors. The combined radiomics features based on MRI (including ADC and T2W) and ^{18}F -DCFPyL (2-(3-(1-carboxy-5-[(6- ^{18}F -fluoro-pyridine-3-carbonyl)-amino]-pentyl)-ureido)-pentanedioic acid) PET performed better than did the baseline clinical model. However, the final radiomics model did not significantly benefit from the addition of clinical data in terms of improved predictive value (47). Solari *et al.* investigated the performance of radiomics based on ^{68}Ga -PSMA-11 PET/MR to predict postoperative ISUP grade. According to their research, the radiomics model combining ADC and PET features had the highest accuracy (accuracy = 0.82). Overall, the ADC exhibited promising outcomes in single-modality models, and its accuracy improved with the incorporation of T2W- or PET-derived features (48). Furthermore, Feliciani *et al.* developed radiomics models based on MRI-ADC and ^{68}Ga -PSMA-11 PET images to assess their ability to differentiate low-risk patients with PCa (ISUP = 1) from higher-risk patients (ISUP > 1). They found that for predicting patients with low ISUP Group Grade, the ADC and ^{68}Ga -PSMA-11 PET radiomics feature-based models were equal and complementary (49). Overall, there is limited research on combining ^{18}F -PSMA-1007 PET and MRI radiomics in PCa. The majority of radiomics studies of combined PET and MRI are based on ^{68}Ga -labeled PSMA tracers and may not be exactly comparable to our ^{18}F -PSMA-1007 tracer as they have different resolutions and uptake patterns. Furthermore, most studies have focused on predicting ISUP Group Grade, and there is limited literature on early persistent PSA after RP.

As stated above, our finding may have implications for clinical practice. Our combined model outperformed both the radiomics models and the clinical baseline model in assessing early persistent PSA. This model was constructed from clinical variables (age + initial PSA) and PET- and mpMRI- (including T2W, DWI, and

ADC) derived radiomics features. The radiomics features included first-order features such as “lbp-3D-m2_first-order_90Percentile” for DWI, “lbp-3D-k_first-order_Median” for T2W, and “wavelet-LLL_first-order_Minimum” for ADC. Here, “lbp” (local binary patterns) is a texture analysis method that captures local spatial variations in voxel intensity. The “m2” and “k” in “lbp-3D-m2” and “lbp-3D-k” refer to different lbp radius and neighborhood configurations, which influence the granularity of the extracted texture features. The “90Percentile” represents the 90th percentile of intensity values in the transformed image, while the “Median” denotes the median intensity value. “Wavelet-LLL” refers to low-low-low (LLL) frequency wavelet decomposition, which applies low-pass filtering along three orthogonal axes (x, y, z) to preserve coarse structural information while removing high-frequency noise. “Minimum” corresponds to the minimum voxel intensity value after wavelet transformation. These features quantitatively describe the distribution and texture characteristics of the medical images, aiding in the differentiation of tissue properties. Additionally, it was also discovered that persistent PSA could be predicted through the use of a few second-order features including “GLZLM_ZLNU” for PET, which quantifies the nonuniformity of the gray levels or the length of the homogeneous zones. Specifically, GLZLM stands for Gray Level Zone Length Matrix, a method that captures the spatial distribution of gray levels in an image. The ZLNU (Zone Length Nonuniformity) measures the variability in the length of contiguous zones with the same gray level intensity. A higher ZLNU value indicates greater heterogeneity in zone lengths, providing information about the complexity and texture of the image. Other second-order features such as GLSZM, GLRLM, and GLCM, reflect the spatial distribution of gray levels. In summary, each of these radiomics features characterized tumor heterogeneity and served a complementary role in predicting persistent PSA. It should be mentioned that while our study primarily focused on evaluating the predictive ability of preoperative ¹⁸F-PSMA-1007 PET- and mpMRI-derived radiomics features in predicting PSA persistence after RP, we acknowledge that preoperative molecular staging, such as PSMA PET findings related to lymph node involvement, could potentially enhance the model’s predictive accuracy. Due to our study population consisting of patients with localized PCa and the small number of patients with preoperative nodal involvement on imaging (n=6), the molecular stage was not included as a variable in the clinical

model. In clinical practice, early PSA monitoring following RP can help identify high-risk patients with worse oncologic outcomes. Therefore, our combined model may improve clinical treatment. Our study roughly evaluated the value of radiomics features in predicting early postoperative PSA. Further prospective studies are required to assess whether using PET and mpMRI radiomics in treatment selection results in better patient outcomes.

There are several limitations in our study. First, potential selection bias could have arisen due to the study’s retrospective design. Second, persistent PSA is a relatively rare event, and we thus only examined a limited number of events in our study. This could have led to the overfitting of models, potentially constraining the models’ generalizability. The DeLong test indicated no statistically significant difference in the AUC values between the combined model and radiomics models. The small patient population and the imbalanced dataset may partly explain this. The statistical power of the findings may be limited. Additionally, although we used whole-prostate segmentations in our study to reduce complexity and achieve effective radiomics, our segmentation strategy could still be improved. Future research could be enhanced by implementing a fully automated fine-segmentation approach through the appropriate training of a deep learning-based algorithm (50,51). Finally, it is important to note that our clinical baseline model included only age and initial PSA; however, Hao *et al.* demonstrated that preoperative PSA, high-risk group, preoperative ISUP grades 2–5, percentage of positive cores, and other postoperative pathological indicators are independent risk factors for PSA persistence in patients with PCa following RP (52). Due to limitations in data collection and insufficient documentation, biopsy ISUP Group Grade was excluded from our baseline model analysis, which might have affected the predictive performance of the clinical model. Future studies will be focused on the gathering of more extensive and comprehensive multicenter data to address the abovementioned limitations.

Conclusions

In our study, we demonstrated the potential of radiomics based on preoperative images to predict persistent PSA after RP. We successfully developed and validated a novel, combined radiomics-clinical model that outperformed all the other radiomic models and clinical baseline model in predicting postoperative persistent PSA. This highlights the value of the radiomics features derived from

¹⁸F-PSMA-1007 PET and mpMRI in complementing clinical variables for early postoperative risk stratification in PCa. Further verification studies are needed to confirm the reproducibility and clinical utility of this method.

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Footnote

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Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. This study was conducted in accordance with the Declaration of Helsinki (as revised in 2013) and was approved by the Ethics Committee of The First Affiliated Hospital of Wenzhou Medical University (No. KY2022-R012). Informed consent was waived due to the retrospective nature of the analysis.

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